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RAW SEQUENCE LISTING
PATENT APPLICATION: US/09/970,088

DATE: 06/14/2002
TIME: 15:53:33

Input Set : A:\71415062.app
Output Set: N:\CRF3\06142002\I970088.raw

<110> APPLICANT: GRAVEREAUX, EDWIN C.
SILVER, MARCY
ISNER, JEFFREY M.
YOON, YOUNG-SUP
<120> TITLE OF INVENTION: USE OF LYMPHANGIOGENIC AGENTS TO TREAT LYMPHATIC DISORDERS
<130> FILE REFERENCE: 71417/55062
<140> CURRENT APPLICATION NUMBER: 09/970,088
<141> CURRENT FILING DATE: 2001-10-02
<150> PRIOR APPLICATION NUMBER: 60/237,171
<151> PRIOR FILING DATE: 2000-10-02
<160> NUMBER OF SEQ ID NOS: 14
<170> SOFTWARE: PatentIn Ver. 2.1
<210> SEQ ID NO: 1
<211> LENGTH: 8
<212> TYPE: PRT
<213> ORGANISM: Artificial Sequence
<220> FEATURE:
<223> OTHER INFORMATION: Description of Artificial Sequence: Illustrative peptide
<400> SEQUENCE: 1
Asn Val Ser Asp Ser Leu Glu Met
1 5
<210> SEQ ID NO: 2
<211> LENGTH: 7
<212> TYPE: PRT
<213> ORGANISM: Artificial Sequence
<220> FEATURE:
<223> OTHER INFORMATION: Description of Artificial Sequence: Illustrative peptide
<400> SEQUENCE: 2
Trp Glu Phe Pro Arg Glu Arg
1 5
<210> SEQ ID NO: 3
<211> LENGTH: 24
<212> TYPE: DNA
<213> ORGANISM: Artificial Sequence
<220> FEATURE:
<223> OTHER INFORMATION: Description of Artificial Sequence: Synthetic oligonucleotide
<220> FEATURE:
<221> NAME/KEY: modified_base
<222> LOCATION: (18)

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```
     63 <223> OTHER INFORMATION: A, T, C or G
     65 <400> SEQUENCE: 3
W--> 66 aacgtgagyg actcsytnga ratg                                        24
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     70 <211> LENGTH: 21
     71 <212> TYPE: DNA
     72 <213> ORGANISM: Artificial Sequence
     74 <220> FEATURE:
     75 <223> OTHER INFORMATION: Description of Artificial Sequence: Synthetic
     76       oligonucleotide
     78 <400> SEQUENCE: 4
     79 cckytcyckg ggraaytccc a                                           21
     82 <210> SEQ ID NO: 5
     83 <211> LENGTH: 21
     84 <212> TYPE: DNA
     85 <213> ORGANISM: Artificial Sequence
     87 <220> FEATURE:
     88 <223> OTHER INFORMATION: Description of Artificial Sequence: Primer
     90 <400> SEQUENCE: 5
     91 tatggtacaa agatgagagg c                                           21
     94 <210> SEQ ID NO: 6
     95 <211> LENGTH: 21
     96 <212> TYPE: DNA
     97 <213> ORGANISM: Artificial Sequence
     99 <220> FEATURE:
    100 <223> OTHER INFORMATION: Description of Artificial Sequence: Primer
    102 <400> SEQUENCE: 6
    103 acaggtattc acattgctcc t                                           21
    106 <210> SEQ ID NO: 7
    107 <211> LENGTH: 420
    108 <212> TYPE: DNA
    109 <213> ORGANISM: Oryctolagus cuniculus
    111 <400> SEQUENCE: 7
    112 cggtgcgcgg tggccggggc acacgtgccc agcatcgtat ggtacaaaga tgagaggctg 60
    113 ctgcaagaag aatctggaat cgacctcgcg gactcgaacc agaggctgag catccagcgc 120
    114 gtgcgcgagg aggacgcggg ccgctatctg tgcagcgtgt gcaacgccaa gggctgcgtc 180
    115 aactcctccg ccagcgtagc tgtgggaggc gccgaagata gaggcagcat ggagatcgtg 240
    116 atcctcgtgg gcaccggcgt cattgccgtg ttcttttggg tcctcctcct gctcatcttc 300
    117 tgtaacatga ggaggccagc ccacgcggac atcaagacgg gctacttgtc catcatcatg 360
    118 gatcccgggg aggtgcctct ggaggagcaa tgtgaatacc tgtcctacga cgccagccag 420
    121 <210> SEQ ID NO: 8
    122 <211> LENGTH: 420
    123 <212> TYPE: DNA
    124 <213> ORGANISM: Bos sp.
    126 <400> SEQUENCE: 8
    127 cggtgcccag tggctgggac gcacgtaccc agcatcgtgt ggtacaaaga tgagaagctg 60
    128 ctggaagaag agtccggaat cgacctggcg gactcgaacc agaggctgag catccagcgc 120
    129 gtgcgcgagg aggacgcggg ccactatctg tgcagtgtgt gcaacgccaa gggctgtgtc 180
    130 aactcctctg ccagcgtggc tgtggaaggc tctgaggata aaggcagcat ggagatcgtg 240
```

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131 atccttgttg gcaccggagt catcgctgtc tttttctggg tcctccttct cctcatcttc 300
132 tgtaacatga ggaggccaac ccatgcagac atcaagactg gctacttgtc catcatcatg 360
133 gaccccgggg aggtgccttt ggaggagcag tgtgaatacc tgtcctacga tgctagtcaa 420
136 <210> SEQ ID NO: 9
137 <211> LENGTH: 420
138 <212> TYPE: DNA
139 <213> ORGANISM: Homo sapiens
141 <400> SEQUENCE: 9
142 cagtgcttgg tggccggagc gcacgcgccc agcatcgtgt ggtacaaaga cgagaggctg 60
143 ctggaggaaa agtctggagt cgacttggcg gactccaacc agaagctgag catccagcgc 120
144 gtgcgcgagg aggatgcggg acgctatctg tgcagcgtgt gcaacgccaa gggctgcgtc 180
145 aactcctccg ccagcgtggc cgtggaaggc tccgaggata agggcagcat ggagatcgtg 240
146 atccttgtcg gtaccggcgt catcgctgtc ttcttctggg tcctcctcct cctcatcttc 300
147 tgtaacatga ggaggccggc ccacgcagac atcaagacgg gctacctgtc catcatcatg 360
148 gaccccgggg aggtgcctct ggaggagcaa tgcgaatacc tgtcctacga tgccagccag 420
151 <210> SEQ ID NO: 10
152 <211> LENGTH: 420
153 <212> TYPE: DNA
154 <213> ORGANISM: Mus sp.
156 <400> SEQUENCE: 10
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158 ctggagaaag agtcgggaat cgacctggca gactccaatc agaggctgag catccagcgc 120
159 gtgcgcgagg aggacgcagg tcgttatctg tgcagcgtgt gcaatgccaa gggctgcgta 180
160 aactcctctg ccagcgtggc agtggaaggc tctgaagata aaggcagcat ggagattgtg 240
161 atactcattg gcactggcgt catcgcagtt ttcttctggg tcctcctcct gctcatcttc 300
162 tgtaacatga aaaggcctgc ccatgcagac atcaagacgg gctacctgtc catcatcatg 360
163 gaccccgggg aggtgccttt ggaggagcag tgtgaatacc tgtcctatga cgccagccag 420
166 <210> SEQ ID NO: 11
167 <211> LENGTH: 140
168 <212> TYPE: PRT
169 <213> ORGANISM: Oryctolagus cuniculus
171 <400> SEQUENCE: 11
172 Arg Cys Ala Val Ala Gly Ala His Val Pro Ser Ile Val Trp Tyr Lys
173   1               5                  10                  15
175 Asp Glu Arg Leu Leu Gln Glu Glu Ser Gly Ile Asp Leu Ala Asp Ser
176              20                  25                  30
178 Asn Gln Arg Leu Ser Ile Gln Arg Val Arg Glu Glu Asp Ala Gly Arg
179          35                  40                  45
181 Tyr Leu Cys Ser Val Cys Asn Ala Lys Gly Cys Val Asn Ser Ser Ala
182      50                  55                  60
184 Ser Val Ala Val Gly Gly Ala Glu Asp Arg Gly Ser Met Glu Ile Val
185  65                  70                  75                  80
187 Ile Leu Val Gly Thr Gly Val Ile Ala Val Phe Phe Trp Tyr Leu Leu
188                  85                  90                  95
190 Leu Leu Ile Phe Cys Asn Met Arg Arg Pro Ala His Ala Asp Ile Lys
191             100                 105                 110
193 Thr Gly Tyr Leu Ser Ile Ile Met Asp Pro Gly Glu Val Pro Leu Glu
194         115                 120                 125
196 Glu Gln Cys Glu Tyr Leu Ser Tyr Asp Ala Ser Gln
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Input Set : A:\71415062.app
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197     130                 135                 140
200 <210> SEQ ID NO: 12
201 <211> LENGTH: 140
202 <212> TYPE: PRT
203 <213> ORGANISM: Bos sp.
205 <400> SEQUENCE: 12
206 Arg Cys Pro Val Ala Gly Thr His Val Pro Ser Ile Val Trp Tyr Lys
207   1               5                  10                  15
209 Asp Glu Lys Leu Leu Glu Glu Glu Ser Gly Ile Asp Leu Ala Asp Ser
210              20                  25                  30
212 Asn Gln Arg Leu Ser Ile Gln Arg Val Arg Glu Glu Asp Ala Gly His
213          35                  40                  45
215 Tyr Leu Cys Ser Val Cys Asn Ala Lys Gly Cys Val Asn Ser Ser Ala
216      50                  55                  60
218 Ser Val Ala Val Glu Gly Ser Glu Asp Lys Gly Ser Met Glu Ile Val
219  65                  70                  75                  80
221 Ile Leu Val Gly Thr Gly Val Ile Ala Val Phe Phe Trp Tyr Leu Leu
222                  85                  90                  95
224 Leu Leu Ile Phe Cys Asn Met Arg Arg Pro Thr His Ala Asp Ile Lys
225             100                 105                 110
227 Thr Gly Tyr Leu Ser Ile Ile Met Asp Pro Gly Glu Val Pro Leu Glu
228         115                 120                 125
230 Glu Gln Cys Glu Val Leu Ser Tyr Asp Ala Ser Gln
231     130                 135                 140
234 <210> SEQ ID NO: 13
235 <211> LENGTH: 140
236 <212> TYPE: PRT
237 <213> ORGANISM: Homo sapiens
239 <400> SEQUENCE: 13
240 Gln Cys Leu Val Ala Gly Ala His Ala Pro Ser Ile Val Trp Tyr Lys
241   1               5                  10                  15
243 Asp Glu Arg Leu Leu Glu Glu Lys Ser Gly Val Asp Leu Ala Asp Ser
244              20                  25                  30
246 Asn Gln Lys Leu Ser Ile Gln Arg Val Arg Glu Glu Asp Ala Gly Arg
247          35                  40                  45
249 Tyr Leu Cys Ser Val Cys Asn Ala Lys Gly Cys Val Asn Ser Ser Ala
250      50                  55                  60
252 Ser Val Ala Val Glu Gly Ser Glu Asp Lys Gly Ser Met Glu Ile Val
253  65                  70                  75                  80
255 Ile Leu Val Gly Thr Gly Val Ile Ala Val Phe Phe Trp Val Leu Leu
256                  85                  90                  95
258 Leu Leu Ile Phe Cys Asn Met Arg Arg Pro Ala His Ala Asp Ile Lys
259             100                 105                 110
261 Thr Gly Tyr Leu Ser Ile Ile Met Asp Pro Gly Glu Val Pro Leu Glu
262         115                 120                 125
264 Glu Gln Cys Glu Val Leu Ser Tyr Asp Ala Ser Gln
265     130                 135                 140
268 <210> SEQ ID NO: 14
269 <211> LENGTH: 140
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<212> TYPE: PRT
<213> ORGANISM: Mus sp.
<400> SEQUENCE: 14

Arg	Cys	Pro	Val	Ala	Gly	Ala	His	Val	Pro	Ser	Ile	Val	Trp	Tyr	Lys
1				5					10					15	
Asp	Glu	Arg	Leu	Leu	Glu	Lys	Glu	Ser	Gly	Ile	Asp	Leu	Ala	Asp	Ser
			20					25					30		
Asn	Gln	Arg	Leu	Ser	Ile	Gln	Arg	Val	Arg	Glu	Glu	Asp	Ala	Gly	Arg
		35					40					45			
Tyr	Leu	Cys	Ser	Val	Cys	Asn	Ala	Lys	Gly	Cys	Val	Asn	Ser	Ser	Ala
	50					55					60				
Ser	Val	Ala	Val	Glu	Gly	Ser	Glu	Asp	Lys	Gly	Ser	Met	Glu	Ile	Val
65					70					75					80
Ile	Leu	Ile	Gly	Thr	Gly	Val	Ile	Ala	Val	Phe	Phe	Trp	Val	Leu	Leu
				85					90					95	
Leu	Leu	Ile	Phe	Cys	Asn	Met	Lys	Arg	Pro	Ala	His	Ala	Asp	Ile	Lys
			100					105					110		
Thr	Gly	Tyr	Leu	Ser	Ile	Ile	Met	Asp	Pro	Gly	Glu	Val	Pro	Leu	Glu
		115					120					125			
Glu	Gln	Cys	Glu	Tyr	Leu	Ser	Tyr	Asp	Ala	Ser	Gln				
	130					135					140				

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Please Note:
Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

Seq#:3; N Pos. 18

VERIFICATION SUMMARY DATE: 06/14/2002
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L:66 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3 after pos.:0